

IN THE SPECIFICATION

On page 11, please replace the paragraph from line 8 to line 11 with the following amended paragraph:

In general, a preferred zinc finger framework has the structure:

(A) $X_{0-2} C X_{1-5} C X_{9-14} H X_{3-6} H/C$ (SEQ ID NO:34)

where X is any amino acid, and the numbers in subscript indicated the possible numbers of residues represented by X.

On pages 11 and 12, please replace the paragraph from line 12 on page 11 to line 2 on page 12 with the following amended paragraph:

In a preferred aspect of the present invention, zinc finger nucleic acid binding motifs may be represented as motifs having the flowing primary structure:

(B) $X^a C X_{2-4} C X_{2-3} F X^c X X X X L X X H X X X^b H$ - linker (SEQ ID NO: 22)

wherein X (including X^a , X^b and X^c) is any amino acid. X_{2-4} and X_{2-3} refer to the presence of 2 [[or]] to 4, or 2 or 3, amino acids, respectively, and $X X X X L X X H X X$ between X^c and X^b are designated positions -1, 1, 2,3,4,5,6,7,8, and 9. The Cys and His residues, which together coordinate the zinc metal atom, are marked in bold text and are usually invariant, as is the Leu residue at position +4 in the α -helix.